

# STIC Search Report Biotech-Chem Library

#### STIC Database Tracking Number: 147961

TO: Christine Saoud Location: rem/4e81/4c70

**Art Unit: 1647** 

Wednesday, March 16, 2005 Case Serial Number: 10/016177 From: Barb O'Bryen

**Location: Biotech-Chem Library** 

Remsen 1A69

Phone: 571-272-2518

MoB

barbara.obryen@uspto.gov

Search Notes	(E) 2
	ı



GenCore version 5.1.6 Copyright (c) 1993 - 2005 Compugen Ltd.

nucleic search, using sw model OM nucleic

March 16, 2005, 10:24:47; Search time 4 Seconds (without alignments) 3.428 Million cell updates/sec ü

1 aaagttacattttctctgga..........0ggtaaaaaaaaaaaa 2056 USE10-016-177A-351-COPY 5 Perfect score: Sequence:

Scoring table:

309 189 369 249

69

309 489 369

429

429 609 489 699 549 729

549

789 699 849 729 909 789 696 849

609

IDENTITY NUC Gapop 10.0 , Gapext 0.5

Total number of hits satisfying chosen parameters:

2 seqs, 3335 residues

Searched:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

pubna.seg:\* Database : Pred. No is the number of results predicted by chance to have a core greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

#### SUMMARIES

		Description	Sequence 1, Appli	Sequence 18, Appl	Sequence 1, Appli	Sequence 18, Appl
		ID	US-10-358-281-1	US_09-746-359A-18-COPY	US-10-358-281-1	US-09-746-359A-18-COPY
		<u>B</u>	-	-	Ч	4
		Match Length DB ID	1953	1382	1953	1382
*	Query	Match	93.3	62.1	1.1	1.0
		No. Score	1917.4 93.3 195	1277.8	21.8 1.1	19.6
	1	0	1	7	ო	4
	Result	z			υ	υ

### ALIGNMENTS

US-10-358-281-1

Sequence 1. Application US/10358281

Publication No. US2003017577841

GENERAL INFORMATION:

TITLE OF INVENTION: Interferon Receptor HKAEF92

FILE REPERENCE: PR65P101

CURRENT APPLICATION NUMBER: US/10/358,281

CURRENT PILING DATE: 2003-02-05

PRIOR APPLICATION NUMBER: 09/453,569

PRIOR PILING DATE: 1999-10-02

PRIOR FILING DATE: 1998-06-03

PRIOR FILING DATE: 1998-06-05

NUMBER OF SEQ ID NOS: 2

SOFTWARE: PRECENTING VOINGER: 1988-06-05

NUMBER OF SEQ ID NOS: 2

SEQ ID NO I TYPE: DNA ORGANISM: Homo sapiens FEATURE: NAME/KEY: CDS

1029

909

1090 CCCCAGAAGTTAATCAGCTGCAGAAGGAGGAGGTGGATGCCTGTGCCACGGCTGTGATG 1149 910 ACAGANTGTGTGGAGGTGCAAGGAGGCCATTCCCCTGGTACTGGCCCTGTTTGCCTTT
790 ACAGAATGTGTGGAGTGCAAGGAAAGGCCATTCCCCTGGTAGGCCCTGTTTGCCTTT
790 ACAGAATGTGTGGAGTGCAAGGAAAGGCCATTCCCCTGGTAGGCCTTTGCCCTTG 310 GGAGAAACAGTGTACTATTCTGTCGAATACCAGGGGGAGTACGAGAGCCTGTACACGAGC 430 GATGACATCACGGCCACTGTGCCATACAACCTTCGTGTCAGGGCCACATTGGGGTCACAG 610 CCCCCAGTTTGAGTTCCTTGTGGCCTACTGGAGGAGGAGGAGGCCTGGTGCCGAGGAAGATGTC 430 GGAGAAACAGTGTACTATTCTGTCGAATACCAGGGGGAGTACGAGAGCCTGTACACGAGC 490 CACATCTGGATCCCCAGCAGCTGGTGCTCACTCACTGAAGGTCCTGAGTGTGATGTCACT 550 GATGACATCACGGCCACTGTGCCATACAACCTTCGTGTCAGGGCCACATTGGGCTCACAG 610 ACCICAGCCIGGAGCAICCIGAAGCAICCCITIAAIAGAAACICAACCAICCITACCCGA 490 ACCTCAGCCTGGAGCATCCTGAAGCATCCCTTTAATAGAAACTCAACCATCCTTACCCGA 670 CCTGGGATGGAGATCACCAAAGATGGCTTCCACCTGGTTATTGAGGTGGAGGACCTGGGG 730 CCCCAGITIGAGITICCTIGIGGCCIACIGGAGGAGGAGCCIGGIGCCGAGGAACAIGIC 790 AAAATGGTGAGGAGTGGGGGTATTCCAGTGCACCTAGAAACCATGGAGCCAGGGGCTGCA 250 AIGCAGACTITICACAAIGGITICIAGAAGAAAICIGGACAAGICITITICAIGIGGITITITIC 370 CTCTCTGTACTCTCAACCAACATGAAGCATCTCTTGATGTGGAGCCCAGTGATCGCGCCT Length 1953; 6; Indels DB 1; Query Match 93.3%; Score 1917.4; Best Local Similarity 99.7%; Pred. No. 0; Matches 1921; Conservative 0; Mismatches ; LOCATION: (130)..(1062) US-10-358-281-1 g g g g g 셤 8 g ઠે g ò g à g à g ઠે g ò g à g à g 8 ò ò ò ò ઠે

8 & 8

RESULT 2
US-09-746-159A-18-COPY
US-09-746-159A-18-COPY
Sequence 18, Application US/09746359A
Parent No. US20020042366A1
GENERAL INFORMATION:
APPLICANT: Thompson, Penny

ო

ö

Gaps ö

```
1827 ACTGAAAATGGGATGTGCATGAACACGGAGGATCCATGAACTACTGTAAAGTGTTGACAG 1886
                                                                                                                                                                                                      1887 TGTGTGCACACTGCAGACAGCAGGTGAAATGTATGTGTGCAATGCGACGACGAGAATGCAGAA 1946
                                                                                                                                                                                                                              1807 idcacacaratacarrrcaccreretrecadrerecacacacacacacacacarrracadra 1748
                                                                                           Query Match 1.1%; Score 21.8; DB 1; Length 1953; Best Local Similarity 46.0%; Pred. No. 0; Marches 74; Conservative 0; Mismatches 87; Indels 0;
                                                                                                                                                                                                                                                           1947 GTCAGTAACATGTGCATGTTTGTTGTGCTCCTTTTTTTCTGT 1987
                                                                                                                                                                                                                                                                                1747 GTTCATGGATCCTCCGTGTTCATGCACATCCCATTTTCAGT 1707
                                                                                                                                                                                                                                                                                                                                                     ; Sequence 18, Application US/09746359A
; Patent No. US20020042366A1
; GENERAL INFORMATION:
                         FEATURE:

NAME/KEY: CDS

; LOCATION: (130)..(1062)

US-10-358-281-1
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                              RESULT 4
US-09-746-359A-18-COPY/C
                                                                                                                                                                                                                              셤
                                                                                                                                                   ઠ
                                                                                                                                                                        셤
                                                                                                                                                                                                    ò
                                                                                                                                                                                                                                                        ð
                                                                                                                                                                                                                                                                                 g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   셤
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ઠે
                                                                                                                                                                                                                                                                 1030 CTCCAGTACTCCTGTTGCCCCGTGGTGGTCCTCCCAGACACCTTGAAAATAACCAATTCA 1089
                                                                                                                                                                                                                                                                                                                             1270 TGACCTCTAGACTTTGGGCTTCCACTTGCTTGGCTGAGCAACCCTGGGAAAAGTGACTTC 1329
                                                                                                                                                                                                                   970 GITGGCTTCATGCTGATCCTTGTGGTCGTGCCACTGTTCGTCTGGAAAATGGGCCGGCTG 1029
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1385 TGACCTCTAGACTGGGGGCTGCCACTTGC-TGGCTGAGCAACCCTGGGAAAAGTGACTTC 1443
                      671
                                                    849
                                                                              731
                                                                                                         909
                                                                                                                                   791
                                                                                                                                                                696
                                                                                                                                                                                         851
                                                                                                                                                                                                                                              911
              612 CCCCAGTTTGAGTTCCTTGTGGCCTACTGGACGAGGAGCCTGGTGCCGAGGAACATGTC
                                                   790 AAAATGGTGAGGAGTGGGGGTATTCCAGTGCACCTAGAAACCATGGAGCCAGGGGCTGCA
                                                                    672 AAAATGGTGAGGAGTGTGGGGGGTATTCCAGTGCACCTAGAAACCTTGGAGCCAGGGGCTGCA
                                                                                                         850 TACTGTGTGAAGGCCCCAGACATTCGTGAAGGCCATTGGGAGGTACAGCGCCCTTCAGCCAG
                                                                                                                            732 FACTGTGAAGGCCCAGACATTCGTGAAGGCCATTGGGAGGTACAGCGCCTTCAGCCAG
                                                                                                                                                              910 ACAGAATGTGTGGAGGTGCAAGGAGAGGCCATTCCCCTGGTACTGGCCCTGTTTGCCTTT
                                                                                                                                                                                1444 ATCCCTTCGGTCCTAAGTTTTCTCATCTGTAATGGGGGA 1482
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1330 ATCCCTTCGGTCCCAAGTTTTCTCATCTGTAATGGGGGA 1368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 1. Application US/10358281

Publication No. US20030175778A1

GENERAL INFORMATION

TITLE OF INVENTION: Interferon Receptor HKAEF92

TITLE OF INVENTION: Interferon Receptor HKAEF92

TITLE OF INVENTION: Interferon Receptor HKAEF92

CURRENT PILING DATE: 2003-02-05

PRIOR APPLICATION NUMBER: 09/453,569

PRIOR FILING DATE: 1999-12-02

PRIOR PILING DATE: 1998-06-03

PRIOR PILING DATE: 1998-06-05

NUMBER OF SEQ ID NOS: 2

SOFTWARE: Patentin Ver: 2.0

SEQ ID NO 1

LENGTH: 1953
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 3
US-10-358-281-1/c
                                                                                                                                                                                                                                                                                                                                1090
                g
                                                                                                                                                                                                                                                                                                                                                 6 6 6 6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        В
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              g
ò
                                                   8
                                                                        g
                                                                                                   ઠે
                                                                                                                            g
                                                                                                                                                            ઠે
                                                                                                                                                                                 g
                                                                                                                                                                                                           8 &
                                                                                                                                                                                                                                                                8 &
                                                                                                                                                                                                                                                                                                                      ð
                                                                                                                                                                                                                                                                                                                                                                                                                                                              g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ò
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   g
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ઠે
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ò
```

```
852 CTGTGTGAAGGCCCAGACATTCGTGAAGGCCATTGGGAGGTACAGCGCCTTCAGCCAG 909
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     24; Indels
APPLICANT: Foster, Donald C.
APPLICANT: Thompson, Penny
APPLICANT: Thompson, Penny
APPLICANT: Thompson, Madden, Karen L.
APPLICANT: Madden, Karen L.
APPLICANT: Madden, Karen L.
APPLICANT: Bajan, Maribeth A.
APPLICANT: Bajan, Maribeth A.
APPLICANT: Gapers, Stephen R.
APPLICANT: Chandrasekher, Vasmin A.
APPLICANT: Spot D. NOS: 72
NUMBER OF SCO ID NOS: 72
LENGTH: 1382
LENGTH: LING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 1.0%; Score 19.6; DE
Best Local Similarity 58.6%; Pred. No. 0;
Matches 34; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Search completed: March 16, 2005, 10:24:52 Job time : 5 secs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; FEDS....(1034); LOCATION: (132)...(1034)
US-09-746-359A-18-COPY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Homo sapiens
```

```
March 16, 2005, 10:24:47; Search time 4 Seconds (without alignments) 3.428 Million cell updates/sec
                                                                                                                             GenCore version 5.1.6
Copyright (c) 1993 - 2005 Compugen Ltd.
                                                   OM nucleic - nucleic search, using sw model
                                                                                                                                                                                             Scoring table: IDENTITY_NUC Gapext 0.5
                                                                                                                                     Title:
Perfect score:
Sequence:
                                                                                Run on:
```

Total number of hits satisfying chosen parameters:

2 seqs, 3335 residues Searched:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

pubna.seq:\* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

Description	Sequence 1, Appli Sequence 18, Appli Sequence 1, Appli Sequence 18, Appli			
ID	09-746-359A-18-COPY 10-358-281-1 09-746-359A-18-COPY			
DB	пана			
Query Match Length DB ID	1953 1 005 1382 1 005 1953 1 05 1382 1 05			
Query	93.3 62.1 1.1			
Score	1917.4 93.3 1277.8 62.1 21.8 1.1 19.6 1.0			
Result No.	0 0 4-10 10 4-			

```
JOSEPH CONTRIBUTION OF STATE OF SEQUENCE O
ALIGNMENTS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: CDS
```

```
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                12
                                                                                                                                                                                                                                  310
                                                                                                                                                                                                                                                               252
                                                                                                                                                                                                                                                                          430
                                                                                                                                                                                                                                                                                             490
                                                                                                                                                                                                                                                                                                       372
                                                                                                                                                                                                              250
                                                                                                                                                                                                                                           192
                                                                                                                                                                                                                                                     370
                                                                                                                                                                                                                                                                                   312
                                                                                                                                                                      à
                                                                                                                                                                               셤
                                                                                                                                                                                          ઠે
                                                                                                                                                                                                  ď
                                                                                                                                                                                                              δ
                                                                                                                                                                                                                     g
                                                                                                                                                                                                                              ઠે
                                                                                                                                                                                                                                        요
                                                                                                                                                                                                                                                 ያ ያ
                                                                                                                                                                                                                                                                                                    g
                                                                                                                                                                                                                                                                        ò
                                                                                                                                                                                                                                                                                g
                                                                                                                                                                                                                                                                                           Š
                                                                                                                                                                                                                                                                                                                 ò
                                                                                                                                                                                                                                                                                                                        셤
                                                                                                                                                                                                                                                                                                                                  ሯ
                                                                                                                               CCCCAGAAGTTAATCAGCTGCAGAAGGGAGGAGGTGGATGCCTGTGCCACGGCTGTGATG 1029
                     1089
                               1269
                                   CGCCACGGACAAGGGATGAGAAGTAGGAAGAGCCTGTTGTCTACAAGTCTAGAAGCAA 1329
                                                                       1389
                                                                           1449
                                                                                               1270 TCTAGACTGGGGGCTGCCACTTGCTGGAACACACCTGGGAAAAGTGACTTCATCCCT 1329
                                                                                                             1330 TCGGTCCTAAGTTTTCTCATCTGTAATGGGGGAATTACCTACACACCTGCTAAACACACA 1389
                                                                                                                                                    1030 TCTCCTGAGGAACTCCTCAGGGCCTGGATCTCATAGGTTTGCGGAAGGGCCCAGGTGAAG
                                                        1390 TCTAGACTGGGGGTGCCACTTGCTGGCTGAGCAACCCTGGGAAAAGTGACTTCATCCT
                            CCGAGAACCTGGTCTGCATGACATGGAAACCATGAGGGGACAAGTTGTGTTTTTTT
                                                                                                                                                                                                                                                                                                                                          RESULT 2
US-09-746-359A-18-COPY
Sequence 18, Application US/09746359A
Fatent No. US20020042366A1
GENERAL INFORMATION:
APPLICANT: Thompson, Penny
                                                                                                                                                                                                                                                                                                                           1930 AAAAAA 1936
                                                                                                                                                                                                                                                                                                                  2050 AAAAAA 2056
                            1210
                                      1090
                                                 1270
                                                                     1330
                                                                                                                                                                                                                                    1810
                                                                                                                                                                                                                                              1690
                 a
                           ò
                                    유
                                              ò
                                                        셤
                                                                    ò
                                                                             qq
                                                                                        ò
                                                                                                 g
                                                                                                                                                                                                                                                                                                                  ò
                                                                                                           ò
                                                                                                                    g
                                                                                                                                                                                                                                                                                               8
                                                                                                                             8 8 8
                                                                                                                                                           d
                                                                                                                                                                    8 S
                                                                                                                                                                                                                                                                g
                                                                                                                                                                                                                                                                         ò
                                                                                                                                                                                         장염
                                                                                                                                                                                                             8 %
                                                                                                                                                                                                                                ò
                                                                                                                                                                                                                                           원
                                                                                                                                                                                                                                                     ò
```

249 72 AIGGACAGAAIGCTITAITITIGGAAAGAACAAIGTICIAGGICAAACTGAGTCTACCAA 131 ATGCAGACTTTCACAATGGTTCTAGAAGAAATCTGGACAAGTCTTTTCATGTGGTTTTTTC 309 429 549 CICTCTGTACTCTCAACCAACATGAAGCATCTCTTGATGTGGAGCCCAGTGATCGCGCCT 311 GGAGAAACAGTGTACTATTCTGTCGAATACCAGGGGGAGTACGAGAGCCTGTACACGAGC 489 371 550 GATGACATCACGGCCACTGTGCCATACAACCTTCGTGTCAGGGCCACATTGGGCTCACAG 609 699 729 71 190 AIGGACAGAAIGCITTAITIIGGAAAGAAACAAIGITCIAGGICAAACIGAGICIACCAA 130 GGCTTCGCTGCGACTCAGACCTCCAACATAGCATTCTGAAGAAGATGGCTGAG CCTGGGATGGAGATCACCAAAGATGGCTTCCACCTGGTATTGAGCTGGAGGACCTGGGG GGAGAAACAGTGTACTATTCTGTCGAATACCAGGGGAGTACGAGAGCCTGTACACGAGC CACATCTGGATCCCCAGCAGCTGGTGCTCACTCACTGAAGGTCCTGAGTGTGATCTCACT 610 ACCTCAGCCTGGAGCATCCTGAAGCATCCCTTTAATAGAAACTCAACCATCCTTACCCGA 8 DB 1; Length 1382; Query Match 62.1%; Score 1277.8; DB 1; Length Best Local Similarity 98.5%; Pred. No. 0; Marches 1339; Conservative 0; Mismatches 12; Indels APPLICANT: FOBIER, Donald C.
APPLICANT: Xu, Wenfeng
APPLICANT: Xu, Wenfeng
APPLICANT: Kelly, James D.
APPLICANT: Specher, Cindy A.
APPLICANT: Specher, Cindy A.
APPLICANT: Blumberg, Hal
APPLICANT: Bagan, Maribeth A.
APPLICANT: Chandrasekher, Yasmin A.
APPLICANT: Chandrasekher, Yasmin A.
APPLICANT: No. US20020042366A1ak, Julia E.
APPLICANT: No. US20020042366A1ak, Julia E.
TITLE OF INVENTION: Method for Treating Inflammation; FILE REPERENCE: 99-108
CURRENT PAPLICATION NUMBER: US/09/746,359A
CURRENT PALING DATE: 2001-05-21
PRIOR APPLICATION NUMBER: 60/171,969
PRIOR APPLICATION NUMBER: 60/171,969
PRIOR PLING DATE: 1099-12-23
PRIOR PLING DATE: 2000-06-22
NUMBER OF SEQ ID NOS: 72
SEQ ID NOS: 72
SEQ ID NO 18
LENGTH: 1382
TWUER: NNA ; FEATURE: ; NAME/KEY: CDS ; LOCATION: (132)...(1034) US-09-746-359A-18-COPY 670 d

```
1090 CCCCAGAAGTTAATCAGCTGCAGAAGAGAAGATGGATGCCTGTGCCACGGCTGTGATG 1149
          730 CCCCAGTITGAGTICCTIGGGCCTACTGGAGGAGGGAGCCTGGTGCCGAGGAACAIGIC 789
                       850 TACTGTGAAGGCCCAGACATTCGTGAAGGCCATTGGGAGGTACAGCGCCTTCAGCCAG 909
                                                                                               8 S
                                                                6 6 6 6 6
                                                                                                    8 8 8 8
```

RESULT 3
US-10-358-281-1/C
US-10-358-281-1/C
Sequence 1, Application US/10358281
Fublication No. US2003015778A1
Fublication No. US2003015778A1
FUBLERAL INPORMATION:
FILE REPERENCE: PF665PLT
CURRENT APPLICATION NUMBER: US/10/358,281
CURRENT APPLICATION NUMBER: US/10/358,281
FRIOR PILING DATE: 2003-02-05
FRIOR APPLICATION NUMBER: 09/453,569
FRIOR PLING DATE: 1999-12-02
FRIOR PLING DATE: 1999-12-02
FRIOR PLING DATE: 1999-06-03
FRIOR FILING DATE: 1999-06-03
FRIOR FLING DATE: 1998-06-05
FRIOR FLING DATE: 1998-06-05